



SEQUENCE LISTING

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GUTTERIDGE, ALEX

<120> ADHESION MOLECULES

<130> 674575-2004

<140> 10/615,515

<141> 2003-07-08

<150> PCT/GB02/00107

<151> 2002-01-11

<150> GB 0100750.9

<151> 2001-01-11

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<170> PatentIn Ver. 3.2

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<211> 1284

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Gly Glu Ser Val Ser Ile Pro Thr Glu Asp Ile Ser Glu Pro Met Phe
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His Phe Val Gln Asn Asp Arg Ile Glu Arg Pro Gln Gly Gly Gly Gly
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Gly Ser Gly Ser Gly Gln Gly Gln Ala Ser Gln Asp Gly Glu Gly Gln
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Leu Thr Glu Tyr Lys Thr His Arg Ala Gly Tyr Thr Ala Asn Gly Val
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Pro Ala Asn Ile Ser Val Val Arg Ser Leu Gln Asn Ser Leu Ala Arg
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Arg Thr Ala Met Thr Ala Gly Lys Arg Arg Glu Leu His Ala Leu Glu
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Glu Asn Leu Ala Ile Ile Ser Asn Ser Glu Pro Ala Gln Leu Leu Glu
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Glu Glu Arg Leu Arg Lys Glu Ile Ala Glu Leu Arg Ala Lys Ile Glu
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      245          250          255
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Tyr Ile Leu Leu Tyr Leu Phe Leu Ser Arg Thr Tyr Lys Asn Val Glu
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 Leu Lys Leu Met Asp Glu Val Val Lys Glu Arg Tyr Asn Pro Ala Gln
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 Asp Ser Pro Leu Cys His Glu Ile Leu Ala Lys Lys Leu Leu Pro Val
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 Val Arg Tyr Tyr Ser Tyr Ile Glu Ile Thr Arg Arg Ala His Gln Thr
 370 375 380
 Leu Trp Arg Glu Tyr Glu His Leu Gln Ser Thr Phe Asp Asn Phe Ala
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Pro	Trp	Ala	Glu	Gln	Ala	Arg	Gln	Leu	Val	Asp	Ala	Asn	Ser	Thr	Ile	50	55	60	
Thr	Ser	Ala	Leu	His	Thr	Leu	Phe	Leu	Gln	Arg	Trp	Arg	Leu	Ser	Leu	65	70	75	80
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Leu	Glu	Pro	Ile	Leu	Ala	Asp	Asn	Asn	Thr	Ala	Ala	Gly	Arg	Leu	Trp	115	120	125	
Asp	Met	Ser	Ala	Gly	Gln	Leu	Lys	Arg	Gly	Asp	Tyr	Gln	Leu	Ile	Val	130	135	140	
Lys	Tyr	Gly	Glu	Phe	Leu	Asn	Glu	Gln	Pro	Glu	Leu	Lys	Arg	Leu	Ala	145	150	155	160
Glu	Gln	Leu	Gly	Arg	Ser	Arg	Glu	Ala	Lys	Ser	Ile	Pro	Arg	Asn	Asp	165	170	175	
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Phe	Tyr	Arg	Arg	Leu	Val	Glu	Lys	Gln	Leu	Leu	Thr	Tyr	Arg	Leu	His	225	230	235	240
Gly	Glu	Ser	Trp	Arg	Glu	Lys	Val	Ile	Glu	Arg	Pro	Val	Val	His	Lys	245	250	255	
Asp	Tyr	Asp	Glu	Gln	Pro	Arg	Gly	Pro	Phe	Ile	Val	Cys	Val	Asp	Thr	260	265	270	

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 370 375 380
 Val His Gln His Arg Phe His Ala Val Ala Met Ser Ala His Gly Lys
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 <213> Homo sapiens

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Leu His Arg Phe Leu Cys Phe Gly Ser Glu Gly Gly Thr Tyr Tyr Ile
      35              40              45

Lys Glu Gln Lys Leu Gly Leu Glu Asn Ala Glu Ala Leu Ile Arg Leu
      50              55              60

Ile Glu Asp Gly Arg Gly Cys Glu Val Ile Gln Glu Ile Lys Ser Phe
      65              70              75              80

Ser Gln Glu Gly Arg Thr Thr Lys Gln Glu Pro Met Leu Phe Ala Leu
      85              90              95

Ala Ile Cys Ser Gln Cys Ser Asp Ile Ser Thr Lys Gln Ala Ala Phe
      100             105             110

Lys Ala Val Ser Glu Val Cys Arg Ile Pro Thr His Leu Phe Thr Phe
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Ile Gln Phe Lys Lys Asp Leu Lys Glu Ser Met Lys Cys Gly Met Trp
      130             135             140

Gly Arg Ala Leu Arg Lys Ala Ile Ala Asp Trp Tyr Asn Glu Lys Gly
      145             150             155             160

Gly Met Ala Leu Ala Leu Ala Val Thr Lys Tyr Lys Gln Arg Asn Gly
      165             170             175

Trp Ser His Lys Asp Leu Leu Arg Leu Ser His Leu Lys Pro Ser Ser
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Glu Gly Leu Ala Ile Val Thr Lys Tyr Ile Thr Lys Gly Trp Lys Glu
      195             200             205

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His	Leu	Leu	Thr	Asn	His	Leu	Lys	Ser	Lys	Glu	Val	Trp	Lys	Ala	Leu	260	265	270
Leu	Gln	Glu	Met	Pro	Leu	Thr	Ala	Leu	Leu	Arg	Asn	Leu	Gly	Lys	Met	275	280	285
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Cys	Glu	Lys	Leu	Cys	Asn	Glu	Lys	Leu	Leu	Lys	Lys	Ala	Arg	Ile	His	305	310	315 320
Pro	Phe	His	Ile	Leu	Ile	Ala	Leu	Glu	Thr	Tyr	Lys	Thr	Gly	His	Gly	325	330	335
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Val	Leu	Met	Ala	Met	Ser	Gln	Ile	Pro	Ala	Gly	Gly	Thr	Asp	Cys	Ser	435	440	445
Leu	Pro	Met	Ile	Trp	Ala	Gln	Lys	Thr	Asn	Thr	Pro	Ala	Asp	Val	Phe	450	455	460
Ile	Val	Phe	Thr	Asp	Asn	Glu	Thr	Phe	Ala	Gly	Gly	Val	His	Pro	Ala	465	470	475 480
Ile	Ala	Leu	Arg	Glu	Tyr	Arg	Lys	Lys	Met	Asp	Ile	Pro	Ala	Lys	Leu	485	490	495
Ile	Val	Cys	Gly	Met	Thr	Ser	Asn	Gly	Phe	Thr	Ile	Ala	Asp	Pro	Asp	500	505	510

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<212> PRT

<213> Homo sapiens

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 35 40 45

Ser Tyr Lys Thr Glu Phe Asp Phe Ser Asp Tyr Val Lys Arg Lys Asp
 50 55 60

Pro Asp Ala Leu Leu Lys His Val Lys His Met Leu Leu Leu Thr Asn
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Thr Phe Gly Ala Ile Asn Tyr Val Ala Thr Glu Val Phe Arg Glu Glu
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Leu Gly Ala Arg Pro Asp Ala Thr Lys Val Leu Ile Ile Ile Thr Asp
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Gly Glu Ala Thr Asp Ser Gly Asn Ile Asp Ala Ala Lys Asp Ile Ile
 115 120 125

Arg Tyr Ile Ile Gly Ile Gly Lys His Phe Gln Thr Lys Glu Ser Gln
 130 135 140

Glu Thr Leu His Lys Phe Ala Ser Lys Pro Ala Ser Glu Phe Val Lys
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Lys Lys Ile Tyr Val Ile Glu
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<212> PRT

<213> Homo sapiens

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 Tyr Ala Asn Asn Pro Arg Val Val Phe Asn Leu Asn Thr Tyr Lys Thr
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 Lys Glu Glu Met Ile Val Ala Thr Ser Gln Thr Ser Gln Tyr Gly Gly
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 Asp Leu Thr Asn Thr Phe Gly Ala Ile Gln Tyr Ala Arg Lys Tyr Ala
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 Tyr Ser Ala Ala Ser Gly Gly Arg Arg Ser Ala Thr Lys Val Met Val
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 Val Val Thr Asp Gly Glu Ser His Asp Gly Ser Met Leu Lys Ala Val
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 Ile Asp Gln Cys Asn His Asp Asn Ile Leu Arg Phe Gly Ile Ala Val
 130 135 140
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 Glu Ile Lys Ala Ile Ala Ser Ile Pro Thr Glu Arg Tyr Phe Phe Asn
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 <213> Homo sapiens

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 35 40 45

Phe Arg Ile His Phe Thr Phe Lys Glu Phe Gln Asn Asn Pro Asn Pro
 50 55 60
 Arg Ser Leu Val Lys Pro Ile Thr Gln Leu Leu Gly Arg Thr His Thr
 65 70 75 80
 Ala Thr Gly Ile Arg Lys Val Val Arg Glu Leu Phe Asn Ile Thr Asn
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 Gly Ala Arg Lys Asn Ala Phe Lys Ile Leu Val Val Ile Thr Asp Gly
 100 105 110
 Glu Lys Phe Gly Asp Pro Leu Gly Tyr Glu Asp Val Ile Pro Glu Ala
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 Asp Arg Glu Gly Val Ile Arg Tyr Val Ile Gly Val Gly Asp Ala Phe
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 Arg Ser Glu Lys Ser Arg Gln Glu Leu Asn Thr Ile Ala Ser Lys Pro
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 <212> PRT
 <213> Mus musculus

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 Gln Glu Gly Arg Thr Ala Lys Gln Glu Pro Leu Leu Phe Ala Leu Ala
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 Val Cys Ser Gln Cys Ala Asp Ile Asn Thr Lys Gln Ala Ala Phe Lys
 100 105 110
 Ala Val Pro Glu Val Cys Arg Ile Pro Thr His Leu Phe Thr Phe Ile
 115 120 125

Gln Phe Lys Lys Asp Leu Lys Glu Ser Met Lys Cys Gly Met Trp Gly
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 Arg Ala Leu Arg Lys Ala Val Ala Asp Trp Tyr Asn Glu Lys Gly Gly
 145 150 155 160
 Met Ala Val Ala Leu Val Val Thr Lys Tyr Lys Gln Arg Asn Gly Trp
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 Ser His Lys Asp Leu Leu Arg Leu Ser His Leu Lys Pro Ser Ser Glu
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 Gly Leu Ala Ile Val Thr Lys Tyr Ile Thr Lys Gly Trp Lys Glu Val
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 Gln Glu Met Pro Leu Thr Ala Leu Leu Arg Asn Leu Gly Lys Met Thr
 275 280 285
 Ala Asn Ser Val Leu Glu Pro Gly Asn Ser Glu Val Ser Leu Ile Cys
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 Glu Lys Leu Ser Asn Glu Lys Leu Leu Lys Lys Ala Arg Ile His Pro
 305 310 315 320
 Phe His Val Leu Ile Ala Leu Glu Thr Tyr Arg Ala Gly His Gly Leu
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 Arg Gly Lys Leu Lys Trp Ile Pro Asp Lys Asp Ile Leu Gln Ala Leu
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 Asp Ala Ala Phe Tyr Thr Thr Phe Lys Thr Val Glu Pro Thr Gly Lys
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 Val Phe Thr Asp Asn Glu Thr Phe Ala Gly Gln Val His Pro Ala Val
 465 470 475 480
 Ala Leu Arg Glu Tyr Arg Lys Lys Met Asp Ile Pro Ala Lys Leu Ile
 485 490 495
 Val Cys Gly Met Thr Ser Asn Gly Phe Thr Ile Ala Asp Pro Asp Asp
 500 505 510
 Arg Gly Met Leu Asp Met Cys Gly Phe Asp Thr Ala Ala Leu Asp Val
 515 520 525
 Ile Arg Asn Phe Thr Leu Asp Val Ile
 530 535

<210> 11
 <211> 538
 <212> PRT
 <213> *Xenopus laevis*

<400> 11
 Met Glu Ala Thr Met Asp Gln Thr Gln Pro Leu Asn Glu Lys Gln Val
 1 5 10 15
 Pro Asn Ser Glu Gly Cys Tyr Val Trp Gln Val Ser Asp Met Asn Arg
 20 25 30
 Leu Arg Arg Phe Leu Cys Phe Gly Ser Glu Gly Gly Thr Tyr Tyr Ile
 35 40 45
 Glu Glu Lys Lys Leu Gly Gln Glu Asn Ala Glu Ala Leu Leu Arg Leu
 50 55 60
 Ile Glu Asp Gly Lys Gly Cys Glu Val Val Gln Glu Ile Lys Thr Phe
 65 70 75 80
 Ser Gln Glu Gly Arg Ala Ala Lys Gln Glu Pro Thr Leu Phe Ala Leu
 85 90 95
 Ala Val Cys Ser Gln Cys Ser Asp Ile Lys Thr Lys Gln Ala Ala Phe
 100 105 110
 Arg Ala Val Pro Glu Val Cys Arg Ile Pro Thr His Leu Phe Thr Phe
 115 120 125
 Ile Gln Phe Lys Lys Asp Leu Lys Glu Gly Met Lys Cys Gly Met Trp
 130 135 140

Gly Arg Ala Leu Arg Lys Ala Val Ser Asp Trp Tyr Asn Thr Lys Asp
 145 150 155 160
 Ala Leu Asn Leu Ala Met Ala Val Thr Lys Tyr Lys Gln Arg Asn Gly
 165 170 175
 Trp Ser His Lys Asp Leu Leu Arg Leu Ser His Ile Lys Pro Ala Asn
 180 185 190
 Glu Gly Leu Thr Met Val Ala Lys Tyr Val Ser Lys Gly Trp Lys Glu
 195 200 205
 Val Gln Glu Ala Tyr Lys Glu Lys Glu Leu Ser Pro Glu Thr Glu Lys
 210 215 220
 Val Leu Lys Tyr Leu Glu Ala Thr Glu Arg Val Lys Arg Thr Lys Asp
 225 230 235 240
 Glu Leu Glu Ile Ile His Leu Ile Asp Glu Tyr Arg Leu Val Arg Glu
 245 250 255
 His Leu Leu Thr Ile His Leu Lys Ser Lys Glu Ile Trp Lys Ser Leu
 260 265 270
 Leu Gln Asp Met Pro Leu Thr Ala Leu Leu Arg Asn Leu Gly Lys Met
 275 280 285
 Thr Ala Asp Ser Val Leu Ala Pro Ala Ser Ser Glu Val Ser Ser Val
 290 295 300
 Cys Glu Arg Leu Thr Asn Glu Lys Leu Leu Lys Lys Ala Arg Ile His
 305 310 315 320
 Pro Phe His Ile Leu Val Ala Leu Glu Thr Tyr Lys Lys Gly His Gly
 325 330 335
 Asn Arg Gly Lys Leu Arg Trp Ile Pro Asp Thr Ser Ile Val Glu Ala
 340 345 350
 Leu Asp Asn Ala Phe Tyr Lys Ser Phe Lys Leu Val Glu Pro Thr Gly
 355 360 365
 Lys Arg Phe Leu Leu Ala Ile Asp Val Ser Ala Ser Met Asn Gln Arg
 370 375 380
 Val Leu Gly Ser Ile Leu Asn Ala Ser Val Val Ala Ala Ala Met Cys
 385 390 395 400
 Met Leu Val Ala Arg Thr Glu Lys Asp Ser His Met Val Ala Phe Ser
 405 410 415
 Asp Glu Met Leu Pro Cys Pro Ile Thr Val Asn Met Leu Leu His Glu
 420 425 430
 Val Val Glu Lys Met Ser Asp Ile Thr Met Gly Ser Thr Asp Cys Ala
 435 440 445

Leu Pro Met Leu Trp Ala Gln Lys Thr Asn Thr Ala Ala Asp Ile Phe
 450 455 460
 Ile Val Phe Thr Asp Cys Glu Thr Asn Val Glu Asp Val His Pro Ala
 465 470 475 480
 Thr Ala Leu Lys Gln Tyr Arg Glu Lys Met Gly Ile Pro Ala Lys Leu
 485 490 495
 Ile Val Cys Ala Met Thr Ser Asn Gly Phe Ser Ile Ala Asp Pro Asp
 500 505 510
 Asp Arg Gly Met Leu Asp Ile Cys Gly Phe Asp Ser Gly Ala Leu Asp
 515 520 525
 Val Ile Arg Asn Phe Thr Leu Asp Leu Ile
 530 535

<210> 12
 <211> 643
 <212> PRT
 <213> *Caenorhabditis elegans*

<400> 12
 Met Ala Asp Glu Leu Asn Glu Phe Gln Glu Ala Gly Asn Phe Asn Glu
 1 5 10 15
 Glu Ala Leu Met Arg Leu Ser Asn Val Cys Ala Arg Leu Arg Arg Met
 20 25 30
 Gln Met Leu Glu Ser Asp Val Glu Ile Thr Val Val Asp Gly Glu Leu
 35 40 45
 Lys Arg Val Pro Arg Gln Met Glu Lys Val Lys Asp Gly Gln Val Glu
 50 55 60
 Asn Asn Ala Gly Gly Phe Val Phe Pro Val Ser Asp Glu Thr Gln Val
 65 70 75 80
 Arg Arg Phe Leu Ile Leu Gly Ser Asp Lys Gly Ser Tyr His Gln Ser
 85 90 95
 Ser Glu Lys Ile Thr Ile Asp Asn Ala Gln Arg Ile Ile Lys Ile Ile
 100 105 110
 Glu Gln Gly Asn Gly His Met Val Leu Lys Glu Leu Ala Leu Ile Asn
 115 120 125
 Ala Glu Asn Arg Asn Pro Lys Met Asn Ala Met Ile Phe Thr Leu Ala
 130 135 140
 Ile Cys Ala Arg Ile Ser Thr His Asp Thr Thr Lys Lys Thr Glu Cys
 145 150 155 160
 Pro Met Leu Asn Ala Tyr Ser Asp Tyr Ile Arg Ala Leu His Asp Ser
 165 170 175

Ala	Leu	Asp	Leu	Ile	Pro	Glu	Val	Cys	Arg	Thr	Pro	Thr	His	Leu	Phe	180	185	190
Glu	Phe	Val	Asp	Tyr	Cys	Gln	Thr	Ile	Ser	Glu	Ser	Thr	Lys	Ala	Gly	195	200	205
Gly	Ala	Lys	Ser	Ser	Thr	Gly	Trp	Gly	Arg	Ser	Met	Arg	Asn	Ala	Ile	210	215	220
Ser	Lys	Trp	Tyr	Thr	Thr	Lys	Thr	Thr	Glu	Lys	Leu	Ala	Met	Leu	Leu	225	230	235
Thr	Lys	Tyr	Pro	Gln	Arg	Glu	Gly	Trp	Ser	His	Arg	Asp	Leu	Phe	Arg	245	250	255
Leu	Ala	His	Pro	Asn	Leu	Met	Asp	Ser	Arg	Ser	His	Gly	Gln	Ser	Glu	260	265	270
Asp	Arg	Leu	Glu	Arg	Glu	Gln	Leu	Phe	Arg	Phe	Ala	Val	Lys	Gly	Asp	275	280	285
Leu	Val	Lys	Arg	Lys	Arg	Lys	Met	Ser	Val	Glu	Glu	Val	Ala	Glu	Val	290	295	300
Glu	Lys	Val	Trp	Asp	Lys	Lys	Ala	Leu	Lys	Leu	Pro	Tyr	Thr	Glu	Glu	305	310	315
Gln	Leu	Ile	Lys	Glu	Glu	Gln	Ser	Arg	Ala	Leu	Asn	Leu	Val	Glu	Ala	325	330	335
Tyr	Leu	Lys	Leu	Lys	Asn	Glu	Gln	Ser	Glu	Glu	Val	Ile	Val	Ala	Ala	340	345	350
Ile	Lys	Lys	His	Gly	Leu	Val	Arg	Glu	His	Leu	Pro	Thr	Thr	Ser	Leu	355	360	365
Asn	Ser	Lys	Leu	Val	Trp	Glu	Thr	Leu	Phe	Asp	Val	Ser	Met	Pro	Met	370	375	380
Thr	Ala	Met	Ile	Arg	Asn	Leu	Ala	Lys	Met	Thr	Val	Val	Gly	Ala	Leu	385	390	395
Asp	Glu	Lys	Arg	Val	Asp	Asn	Ile	Val	Lys	Arg	Leu	Thr	Asp	Gln	Glu	405	410	415
Glu	Leu	Arg	Arg	Ser	Arg	Ile	His	Pro	Ile	Asn	Leu	Leu	Thr	Ala	Arg	420	425	430
Ala	Val	Tyr	Ala	Gln	Gly	Arg	Gly	Asp	Lys	Gly	Ser	Leu	Thr	Trp	Glu	435	440	445
Pro	Asn	Gln	Lys	Ile	Cys	Asp	Ala	Leu	Glu	Ala	Gly	Phe	Tyr	Lys	Ala	450	455	460
Phe	Val	Asn	Ala	Pro	Pro	Thr	Gly	Lys	Arg	Tyr	Cys	Leu	Ala	Leu	Asp	465	470	475
																		480

Val	Ser	Gly	Ser	Met	Thr	Ser	Arg	Val	Ser	Ser	Ser	Pro	Leu	Ser	Cys	485	490	495
Arg	Glu	Ala	Ala	Thr	Gly	Met	Ser	Leu	Ile	Asn	Leu	His	Asn	Glu	Ala	500	505	510
Glu	Val	Arg	Cys	Val	Ala	Phe	Cys	Asp	Lys	Leu	Thr	Glu	Leu	Pro	Phe	515	520	525
Thr	Lys	Asp	Trp	Lys	Ile	Gly	Gln	Val	Asn	Asp	Tyr	Val	Asn	Asn	Leu	530	535	540
Asp	Phe	Gly	Arg	Thr	Asp	Cys	Gly	Leu	Pro	Met	Thr	Trp	Ala	Thr	Glu	545	550	555
Asn	Asn	Leu	Lys	Phe	Asp	Val	Phe	Ile	Ile	Tyr	Thr	Asp	Asn	Asp	Thr	565	570	575
Trp	Ala	Gly	Glu	Ile	His	Pro	Phe	Glu	Ala	Ile	Lys	Lys	Tyr	Arg	Glu	580	585	590
Ala	Ser	Gly	Ile	His	Asp	Ala	Lys	Val	Ile	Val	Met	Ala	Met	Gln	Ala	595	600	605
Tyr	Asp	Tyr	Ser	Ile	Ala	Asp	Pro	Ser	Asp	Ala	Gly	Met	Leu	Asp	Ile	610	615	620
Thr	Gly	Phe	Asp	Ser	Ala	Val	Pro	Gln	Ile	Val	His	Glu	Phe	Val	Thr	625	630	635
Gly	Lys	Ile																640